



Taxonomic Paper

First record of *Trissolcus basalis* (Hymenoptera: Scelionidae) parasitizing *Halyomorpha halys* (Hemiptera: Pentatomidae) in the United States

Rammohan R Balusu[‡], Elijah J Talamas^{§,|}, Ted E Cottrell[¶], Michael D Toews[#], Brett R Blaauw[¤], Ashfaq A Sial[¤], David G Buntin[¤], Henry Y Fadamiro[‡], Glynn Tillman[»]

[‡] Department of Entomology and Plant Pathology, Auburn University, Auburn, AL, United States of America

[§] Florida State Collection of Arthropods, Gainesville, United States of America

[|] USDA/SEL, Washington, DC, United States of America

[¶] United States Department of Agriculture, Agricultural Research Service, Southeastern Fruit & Nut Tree Research Laboratory, Byron, GA, United States of America

[#] Department of Entomology, University of Georgia, Tifton, GA, United States of America

[¤] Department of Entomology, University of Georgia, Athens, GA, United States of America

[¤] Department of Entomology, University of Georgia, Griffin, GA, United States of America

[»] United States Department of Agriculture, Agricultural Research Service, Crop Protection & Management Research Laboratory, Tifton, GA, United States of America

Corresponding author: Rammohan R Balusu (balusrr@auburn.edu)

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Abstract

Background

A parasitoid wasp, *Trissolcus basalis* (Wollaston), was recorded parasitizing eggs of the invasive stink bug *Halyomorpha halys* (Stål) in the United States. This is the first record of this species parasitizing fresh and frozen eggs of *H. halys* in the United States.

New information

First record of *Trissolcus basalis* parasitizing *Halyomorpha halys* eggs in the United States.

Keywords

Parasitoid wasp, endoparasitoid, brown marmorated stink bug

Introduction

The brown marmorated stink bug, *Halyomorpha halys* (Stål), 1855 (Hemiptera: Pentatomidae) (BMSB) is a native of China, Taiwan, South Korea and Japan. Unfortunately, this invasive insect pest has spread to the United States (Lee et al. 2013), where it is both an urban nuisance pest (Inkley 2012) and a serious economic pest of orchard, field and vegetable crops (Leskey et al. 2012a, Leskey et al. 2012b, Rice et al. 2014). The first known *H. halys* populations in the United States were reported in 1996 from Allentown, PA (Hoebelke and Carter 2003). It has now been found in 44 states (StopBMSB 2018).

In the south-eastern U.S., populations of *H. halys* are continuing to expand into the Piedmont and Coastal Plains regions of Georgia and Alabama. *Halyomorpha halys* was first detected in Alabama in 2010. One year later, urban pest management professionals began reporting overwintering brown marmorated stink bugs in homes in the metropolitan Atlanta area. Currently, the brown marmorated stink bug threatens peaches, plums, blueberries, apples, wine grapes, kiwifruit, soybean, cotton, pecan and tomatoes in both states. The tree of heaven, *Ailanthus altissima* (Mill.) Swingle, a tree with seed pods that are a favourite non-crop food source for *H. halys*, also occurs in both states.

Presently, 18 species of hymenopteran endoparasitoids in the genera *Anastatus* Motchulsky (Eupelmidae), *Trissolcus* Ashmead, *Telenomus* Haliday and *Gryon* Haliday (Scelionidae) have been reported to parasitize eggs of *H. halys* in the U.S. (Abram et al. 2014, Rice et al. 2014). As the impact of stink bug parasitoids on this pest was unknown in Georgia and Alabama, a survey to examine parasitism and species composition of parasitoids attacking sentinel egg masses of *H. halys* was conducted in 2017 in regions where populations of *H. halys* had become established.

Materials and methods

Laboratory-reared *H. halys* egg masses were laid on knit cloth (97% cotton, 3% spandex). On 24 June 2018, 30 fresh egg masses (≤ 24 h old) were hung as sentinels on tomato plants for 72 h. Some egg masses (≤ 12 h old) were frozen and held at -20°C for 1–4 d. On 18 October, 30 frozen egg masses were hung as sentinels on plants in cotton and soybean for 72 h. In the laboratory, the collected egg masses were held for emergence of adult parasitoids and emergent wasps were identified using the key of Talamas et al. (2015). Voucher specimens of parasitoids are deposited in the Florida State Collection of Arthropods, Gainesville, Florida (FSCA 00090444, FSCA 00090269).

All egg masses were dissected for dead, immature parasitoids. Determination of *T. basalis* immature stages, mainly third instars, prepupae and pupae, were based on descriptions of *T. basalis* immatures in Volkoff and Colazza (1992) and on descriptions of *H. halys* eggs, parasitised by *T. basalis* every 24 h from oviposition to pupation (G. Tillman, unpublished data).

Two dried point-mounted specimens were selected for DNA extraction and mitochondrial cytochrome c oxidase I (COI) fragment sequencing. Specimens were softened in 70% ethanol for two hours, then DNA was extracted using a DNeasy Blood and Tissue Kit (Qiagen). The DNA samples were quantified using a NanoDrop 2000 spectrophotometer (Thermo Scientific). At least 20 ng of genomic DNA was used per PCR. The 5'-COI region was PCR-amplified using the primers LCO1490 and HCO2198 (Folmer et al. 1994). PCRs were performed at 25 μ l volumes using HiFi HotStart DNA Polymerase (Kapa Biosystems). PCR thermocycle conditions were: 1) initial denaturing at 95°C for 2 min followed by 32 cycles of steps 2–4, 2) 98°C for 30 seconds, 3) 50°C for 30 seconds, 4) 72°C for 40 seconds and 5) final extension at 72°C for 7:00 minutes. PCR products were verified by gel electrophoresis and cleaned for sequencing with QIAquick Gel Extraction Kits (Qiagen). Purified PCR products were Sanger-sequenced in both directions using BigDye Terminator v3.1 (Applied Biosystems) chemistry on a SeqStudio Genetic Analyzer (Applied Biosystems). Sequence reads were trimmed and sequence contigs were assembled in Sequencher 5.4.6 (Gene Codes Corporation). COI sequences, generated during this study, were deposited in GenBank ([MK720833](#), [MK720834](#)).

Taxon treatment

Trissolcus basalis (Wollaston) 1858

- a. scientificName: *Trissolcus basalis*; scientificNameID: urn:lsid:biosci.ohio-state.edu:osuc_names:3189; kingdom: Animalia; phylum: Arthropoda; class: Hexapoda; order: Hymenoptera; family: Scelionidae; genus: *Trissolcus*; specificEpithet: *basalis*; country: United States; stateProvince: Alabama; county: Tuscaloosa; locality: Tuscaloosa, Tuscaloosa Co., AL, U.S.A.; decimalLatitude: 33.21; decimalLongitude: -87.57; georeferenceSources: GNIS-USGS; samplingProtocol: reared from egg; eventDate: 06/17/2017; verbatimEventDate: Jun-17-2017; fieldNotes: [USA: AL: Tuscaloosa. Tomato 6-2, ex. fresh BMSB eggs 17-JUN-2017, Coll. Rammohan Balusu]; individualCount: 1; sex: female; lifeStage: adult; catalogNumber: FSCA 00090269; recordedBy: Balusu, R. (Rammohan); identifiedBy: Talamas, E. J. (Elijah Jacob); dateIdentified: 2019; language: en; institutionCode: Florida State Collection of Arthropods, Gainesville, FL (FSCA); collectionCode: Insects; basisOfRecord: PreservedSpecimen; source: <http://hol.osu.edu/spmlInfo.html?id=FSCA%2000090269>; occurrenceID: urn:lsid:biosci.ohio-state.edu:osuc_occurrences:FSCA_00090269
- b. scientificName: *Trissolcus basalis*; scientificNameID: urn:lsid:biosci.ohio-state.edu:osuc_names:3189; kingdom: Animalia; phylum: Arthropoda; class: Hexapoda; order: Hymenoptera; family: Scelionidae; genus: *Trissolcus*; specificEpithet: *basalis*; country: United States; stateProvince: Alabama; county: Tuscaloosa; locality: Tuscaloosa, Tuscaloosa Co., AL, U.S.A.; decimalLatitude: 33.21; decimalLongitude: -87.57; georeferenceSources: GNIS-USGS; samplingProtocol: reared from egg; eventDate: 07/23/2017; verbatimEventDate: Jul-23-2017; fieldNotes: [USA: AL: Tuscaloosa, Tomato 3-6, ex. fresh BMSB eggs 23-JUL-2017, Coll. Rammohan Balusu]; individualCount: 1; sex: female; lifeStage: adult; catalogNumber: FSCA 00090444; recordedBy: Balusu, R. (Rammohan); identifiedBy: Talamas, E. J. (Elijah Jacob); dateIdentified: 2019; language: en; institutionCode: Florida State Collection of Arthropods, Gainesville, FL (FSCA); collectionCode: Insects; basisOfRecord: PreservedSpecimen; source: <http://hol.osu.edu/spmlInfo.html?id=FSCA%2000090444>; occurrenceID: urn:lsid:biosci.ohio-state.edu:osuc_occurrences:FSCA_00090444

Diagnosis

Trissolcus basalis can be identified from Nearctic congeners by the combination of the following characters: vertex without hyperoccipital carina, netrion sulcus incomplete, mesopleuron with episternal foveae shallowly impressed, metapleuron without setation and without well-defined paracoxal sulcus; T2 striate (Figs 1, 2, 3) (Talamas et al. 2015).